SEQUENCE LISTING

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<110> Carreno, Beatriz M.
      Wood, Clive
      Turner, Katherine
      Collins, Marv
      Gray, Gary S.
      Morris, Donna
      O'Hara, Denise
      Hinton, Paul
      Tsurushita, Naoya
<120> ANTIBODIES AGAINST CTLA4 AND USES THEREFOR
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<150> 60/178,473
<151> 2000-01-27
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gactteetee tetggateet tgeageagtt agtteggggt tgtttttta tagetttete 540
cccatcaatt ga
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426

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Val Phe Cys Lys Ala Met His Val Ala Gln Pro Ala Val Val Leu Ala
                              40
Ser Ser Arg Gly Ile Ala Ser Phe Val Cys Glu Tyr Ala Ser Pro Gly
                         55
Lys Ala Thr Glu Val Arg Val Thr Val Leu Arg Gln Ala Asp Ser Gln
Val Thr Glu Val Cys Ala Ala Thr Tyr Met Met Gly Asn Glu Leu Thr
Phe Leu Asp Asp Ser Ile Cys Thr Gly Thr Ser Ser Gly Asn Gln Val
                                 105
                                                     110
Asn Leu Thr Ile Gln Gly Leu Arg Ala Met Asp Thr Gly Leu Tyr Ile
        115
                            120
Cys Lys Val Glu Leu Met Tyr Pro Pro Pro Tyr Tyr Leu Gly Ile Gly
    130
                        135
Asn Gly Ala Gln Ile Tyr Val Ile Asp Pro Glu Pro Cys Pro Asp Ser
145
                    150
Asp Phe Leu Leu Trp Ile Leu Ala Ala Val Ser Ser Gly Leu Phe Phe
                165
                                    170
Tyr Ser Phe Leu Leu Thr Ala Val Ser Leu Ser Lys Met Leu Lys Lys
            180
                                185
                                                     190
Arg Ser Pro Leu Thr Thr Gly Val Tyr Val Lys Met Pro Pro Thr Glu
        195
Pro Glu Cys Glu Lys Gln Phe Gln Pro Tyr Phe Ile Pro Ile Asn
                        215
                                             220
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tgcactgtct ctgggttttc attaaccagc tatggtgtat attgggttcg ccagcctcca 180 ggaaagggtc tggagtggct gggagtaata tagggctggt ggaaccacaa attataattc 240 ggctctatgt ccagactgag catcagcaaa gacaactcca agagccaagt tttcttaaaa 300 atgagcagtc tgcaaactga tgcacagcc atgtactact gtgccagggg ccccccgcac 360 gctatgatga agagagcta tgctatggac tactgggac aaggaacctc agtcatcgtc 420

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Val Leu Ser Gln Val Gln Leu Lys Glu Ser Gly Pro Gly Leu Val Ala
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Pro Ser Gln Ser Leu Ser Ile Thr Cys Thr Val Ser Gly Phe Ser Leu
Thr Ser Tyr Gly Val Tyr Trp Val Arg Gln Pro Pro Gly Lys Gly Leu
     50
                         55
Glu Trp Leu Gly Val Ile Trp Ala Gly Gly Thr Thr Asn Tyr Asn Ser
Ala Leu Met Ser Arg Leu Ser Ile Ser Lys Asp Asn Ser Lys Ser Gln
Val Phe Leu Lys Met Ser Ser Leu Gln Thr Asp Asp Thr Ala Met Tyr
            100
Tyr Cys Ala Arg Gly Pro Pro His Ala Met Met Lys Arg Gly Tyr Ala
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Met Asp Tyr Trp Gly Gln Gly Thr Ser Val Ile Val Ser Ser
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                                            140
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gtcaccatga cctgcagtgc cacctcaagt ataacttaca tgtcctggta ccagcagaag 180
teaggatect cececagaet cetgatttat gacacateca acetggette tggagtecet 240
gttcgcttca gtggcagtgg gtctgggacc tcttactctc tcacaatcag ccgaatggag 300
gctgaagatg ctgccactta ttactgccag cagtggagta gttacccgct cacgttcggt 360
gctgggacca agctggagct gaaa
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Val Ile Leu Ser Arg Gly Gln Asn Val Leu Thr Gln Ser Pro Ala Ile
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Met Pro Ala Ser Pro Gly Glu Lys Val Thr Met Thr Cys Ser Ala Thr
                             40
                                                  45
Ser Ser Ile Thr Tyr Met Ser Trp Tyr Gln Gln Lys Ser Gly Ser Ser
                         55
Pro Arg Leu Leu Ile Tyr Asp Thr Ser Asn Leu Ala Ser Gly Val Pro
Val Arg Phe Ser Gly Ser Gly Ser Gly Thr Ser Tyr Ser Leu Thr Ile
Ser Arg Met Glu Ala Glu Asp Ala Ala Thr Tyr Tyr Cys Gln Gln Trp
Ser Ser Tyr Pro Leu Thr Phe Gly Ala Gly Thr Lys Leu Glu Leu Lys
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catactgtcc agaggagata tccagatgac ccagtctcca tcctccctat ccgcatcgtt 120
ggggacaggg tcaccataac ctgtagtgcc acctcaagta taacttacat gtcctgtatc 180
agcagaagcc aggaaaggct cccaagcttc tgatttatga cacatccaac ctggctctgg 240
ggtacctagc cgcttcagtg gcagtgggtc tgggaccgac tacacactca caatagcagc 300
ctgcagccag aagattttgc cacttattac tgccagcagt ggagtagtta cccctcacgt 360
tcggtggagg gaccaaggtt gagataaaac gtaagtagaa tccaaagtct aga
                                                                   413
<210> 8
<211> 128
<212> PRT
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Val Ile Leu Ser Arg Gly Asp Ile Gln Met Thr Gln Ser Pro Ser Ser
                                 25
Leu Ser Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys Ser Ala Thr
                                                  45
Ser Ser Ile Thr Tyr Met Ser Trp Tyr Gln Gln Lys Pro Gly Lys Ala
Pro Lys Leu Ile Tyr Asp Thr Ser Asn Leu Ala Ser Gly Val Pro
                     70
Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Tyr Thr Leu Thr Ile
Ser Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Trp
Ser Ser Tyr Pro Leu Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
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                            120
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tgtccttgac ttgcactgtc tctgggtttt cattaacctc atatggtgta tattgggttc 180
gccagcctcc aggaaagggt ctggagtggc tgggagtaat atgggctggt ggtaccacaa 240
attataattc ggctctcatg tccagactga caatcagcaa agacacatcc aagaaccaag 300
tttccttaaa actcagcagt gtgactgcag cggacacagc cgtctactac tgtqcccgag 360
geocecegea egetatgatg aagagagget atgetatgga etaetgggga eaaggaacee 420
tagtcacagt ctcctcaggt gagtccttaa aacctctaga
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Met Ala Val Leu Val Leu Phe Leu Cys Leu Val Ala Phe Pro Ser Cys
                                     10
Val Leu Ser Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Lys
                                 25
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Pro Ser Gln Thr Leu Ser Leu Thr Cys Thr Val Ser Gly Phe Ser Leu 35 40 45

Thr Ser Tyr Gly Val Tyr Trp Val Arg Gln Pro Pro Gly Lys Gly Leu 50 55 60

Glu Trp Leu Gly Val Ile Trp Ala Gly Gly Thr Thr Asn Tyr Asn Ser 65 70 75 80

Ala Leu Met Ser Arg Leu Thr Ile Ser Lys Asp Thr Ser Lys Asn Gln 85 90 95

Val Ser Leu Lys Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val Tyr 100 105 110

Tyr Cys Ala Arg Gly Pro Pro His Ala Met Met Lys Arg Gly Tyr Ala 115 120 125

Met Asp Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser 130 135 140